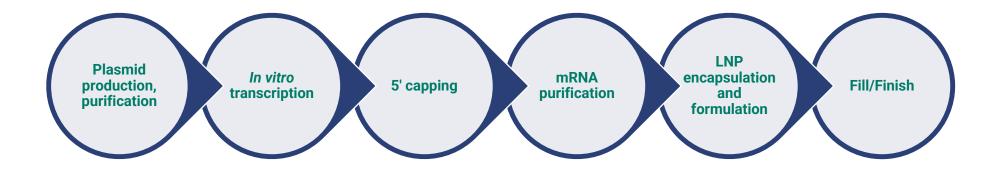
# ° InDevR

ENABLING VACCINES & BIOTHERAPEUTICS **From Bottlenecks to Breakthroughs:** Advancing mRNA Vaccine Development with Same-Day Analytics

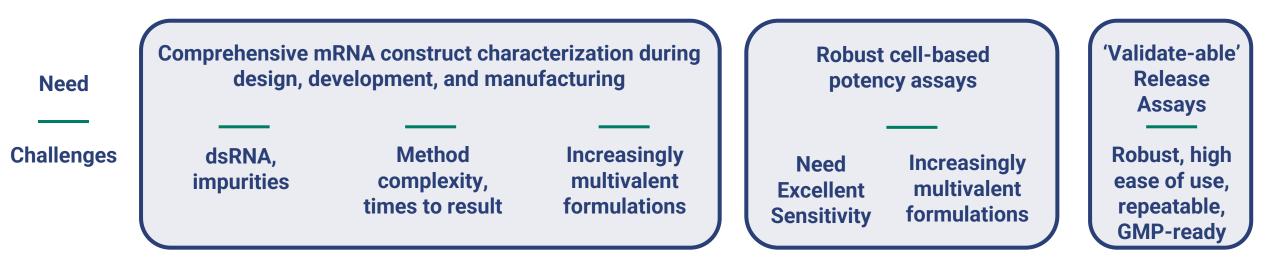
Rachel Gao, PhD, Senior Scientist



# mRNA Analytical Development Bottlenecks



#### **Example Areas of Analytical Technology Constraints**



## InDevR: Next-Generation Vaccine Analytics



Founded in 2003 Boulder, CO

#### InDevR's mission is to **eliminate analytical development bottlenecks** in vaccine development and manufacturing

- 2 decades of streamlining CQA/QC testing for diverse vaccine types (egg-based, cell-based, mRNA)
- **First in industry** to deliver novel strain-responsive off-the-shelf assay kits for **multivalent** influenza antigen testing (HA, NA, NP)
- **Expert Services** division offers tailored solutions for mRNA vaccines as well as traditional vaccines
- Unveiled new cutting-edge facility in Q4 2023 with expanded laboratory capacity

# VaxArray Platform Versatility for mRNA Applications



assessment



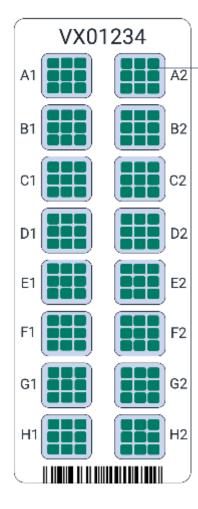


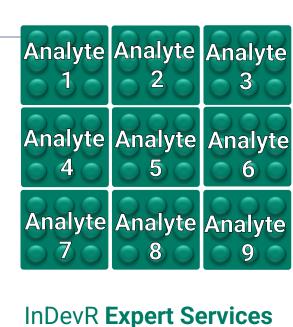
Custom Assay Development Services Available

BioprocessDrug Product TestingClinical/SerologyRapid multiplexed expressed<br/>protein quantitationExtraction-free LNP-mRNA<br/>identity and quantity testingMultiplexed quantification of<br/>antibody levelsNaked mRNA construct identity<br/>and quantity testingRapid multiplexed expressed<br/>protein quantitationRapid multiplexed expressed<br/>protein quantitationRapid 5'-capping efficiencyFinite ConstructClinical/Serology

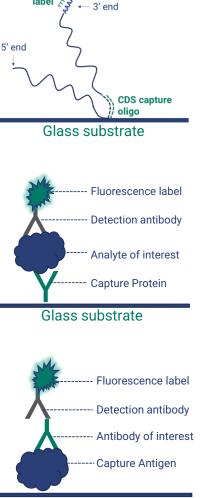
## Single Instrument, Multiple Assay Options

detection label





assays to your needs, multiplexing 50+ analytes



### VaxArray mRNA Oligo Assay

• Measure construct-specific mRNA identity and quantity without extraction

### VaxArray Antibody Assay

 Measure identity and quantity of protein expression from multiplexed mRNA transfection

### VaxArray Antigen Assay

Measure antibody response from patient serological samples

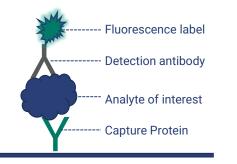
**Glass substrate** 

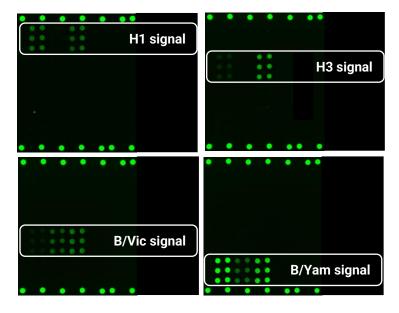


### Applications in mRNA Analytics

# In Vitro Potency: Quantify Proteins Post-Transfection

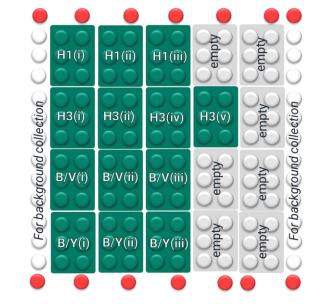
### Assess multivalent expressed proteins with high specificity



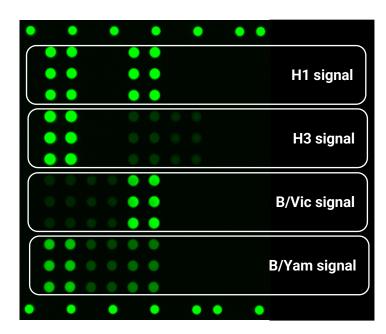


Multiple captures provide reactivity to a broad range of flu strains

Reactivity re-evaluated semi-annually with updated vaccine strains



HA microarray layout

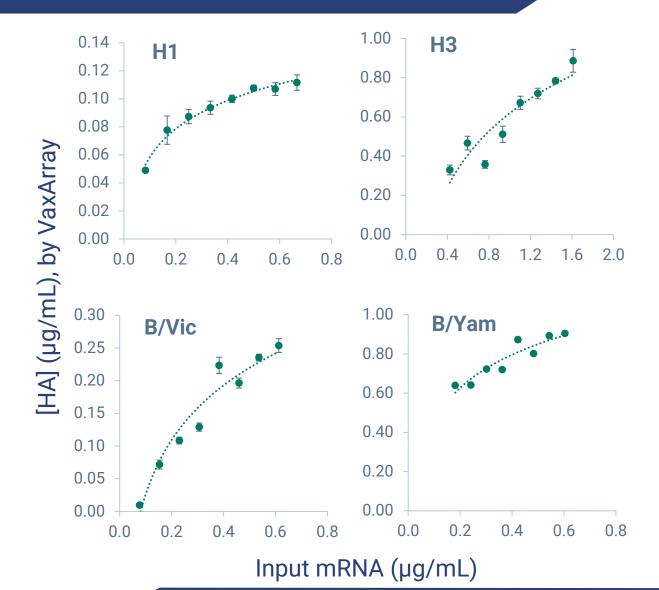


Enables simultaneous quantification of all 4 expressed proteins after quadrivalent mRNA transfection

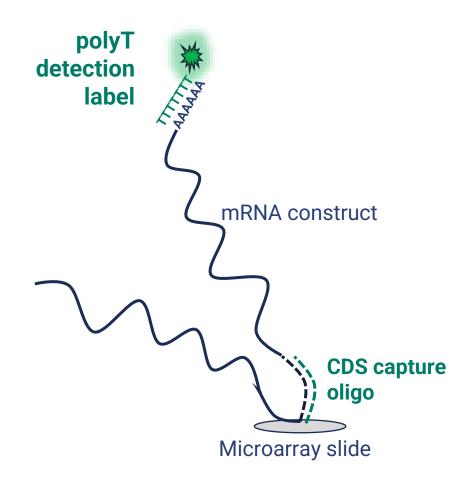
## In Vitro Potency: Quantify Proteins Post-Transfection

#### **Evaluate functional protein during optimization of transfection conditions:**

- Input mRNA concentration
- Sequence optimization
- Transfection time
- Batch to batch
- HEK293-expressed protein post-transfection with a 4-valent mixture of influenza HA mRNA constructs measured by VaxArray HA Assay as a function of input mRNA
- As expected, the different proteins exhibit different expression levels



# Identity and Quantity of mRNA Constructs



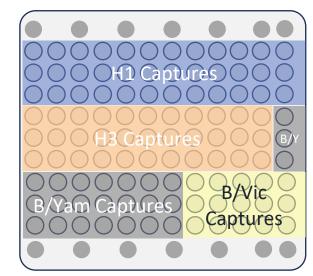
- Identity and quantity of each construct in multivalent mRNA vaccines
- <90-minute time to result
- Capture oligos are designed to target coding regions (CDS) for each construct (ex: each component in a flu vaccine)
- Single detection label for all constructs (custom labels can be designed)
- Works with both naked and LNPencapsulated mRNA (quick 10-min lysis)
- Can be designed on a custom basis

## Identity and Quantity for mRNA Constructs

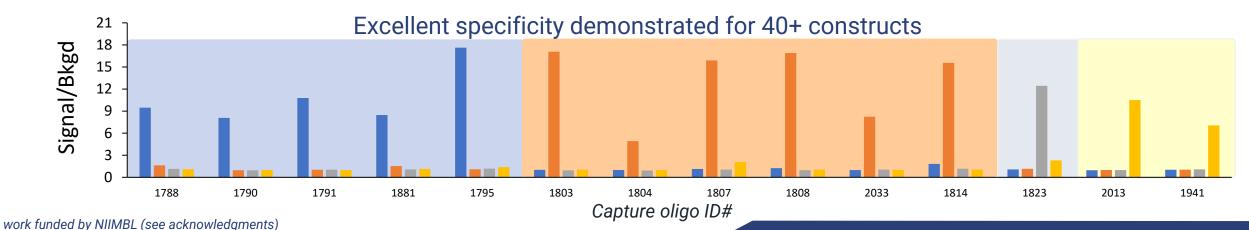
**Case Study:** Assay for Identity and quantity in 4-valent flu HA mRNA vaccines

 Oligos designed to capture regardless of strain and codon optimization scheme in a single assay **32 capture oligos, with oligos specific for each vaccine component** 

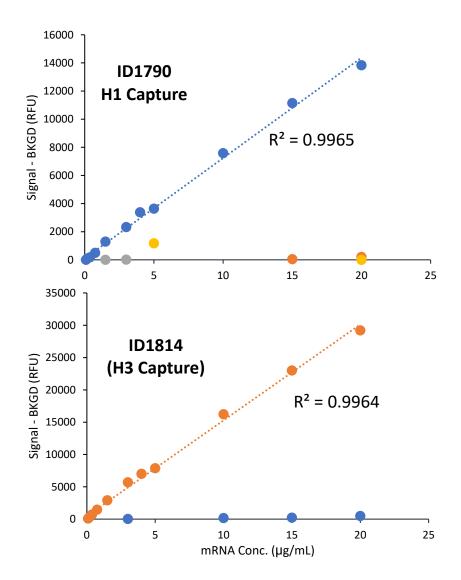
- Multiple strains over time spanning 11 vaccine seasons
- 3 unique suppliers (unique UTRs)
- 4 unique codon optimization methods

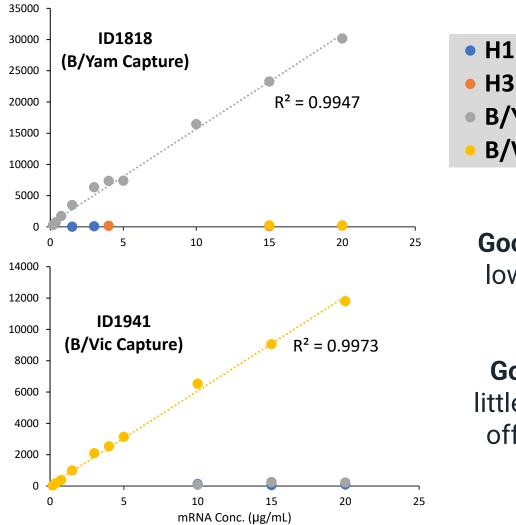


**Prototype Array Layout** 



## Identity and Quantity for mRNA Constructs



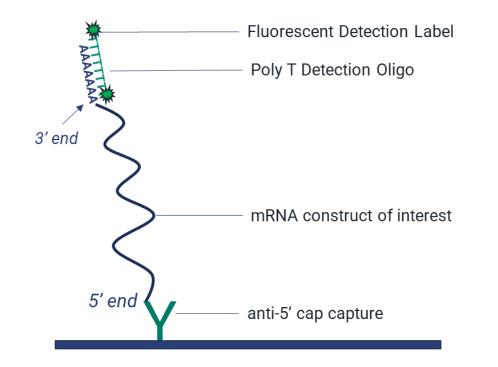


• H3 • B/Y B/V

> Good linearity over low µg/mL mRNA range

**Good specificity**: little to no signal on off-target capture oligos

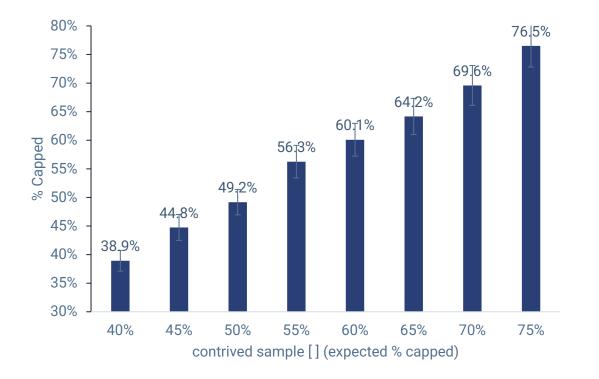
## Rapid 5' Capping Assay for Bioprocess Dev't



- Capture/detection method is universal to any mRNA construct as long as it has a 5' cap and a 3' polyA tail
- < 2-hour time to result
- Assessment of relative capping or quantitative capping efficiency
- For quantitative analysis, user provides characterized matched standard (same construct) if quantification is desired
- Requires an orthogonal measurement of total mRNA on the unknown sample by UV-Vis or other method - therefore material must be reasonably pure

# Assay Can Differentiate Wide Range of % Capped

- Contrived samples created by mixing uncapped and capped GFP mRNA construct
- Samples differed by 5% capping; run alongside a dilution series of known capped standard



Expected Value	Measured Capping Efficiency (95% CI)	Difference from Expected
40%	38.9 (38.4 - 39.4)%	-1.1%
45%	44.8 (44.1 – 45.3)%	-0.2%
50%	49.2 (48.5 - 49.8)%	-0.8%
55%	56.3 (55.5 – 57.0)%	+1.3%
60%	60.1 (59.3 - 60.8)%	+0.1%
65%	64.2 (63.3 - 65.0)%	-0.8%
70%	69.6 (68.7 - 70.5%	-0.4%
75%	76.5 (75.5% - 77.5)%	+1.5%

- All samples statistically differentiated (p-value <0.05 and non-overlapping confidence intervals)
- 8/8 samples were less than 1.5% difference from expected



Funding (Universal influenza HA mRNA assay): This work was performed under a Project Award Agreement from the National Institute for Innovation in Manufacturing Biopharmaceuticals (NIIMBL) and financial assistance award 70NANB17H002 from the U.S. Department of Commerce, National Institute of Standards and Technology.

#### Collaborators/Materials Providers

- Dr. Norbert Pardi, University of Pennsylvania, collaborator on NIIMBL funded project
- 5' capping assay, materials provided and used during development:
  - North Carolina State University/BTEC, Dr. Jennifer Pancorbo
  - Cisterna, Dr. Hari Bhaskaran
  - Aldevron

InDevR's Expert Services and R&D Teams

# We Want to Hear From You



### Scan to connect with InDevR's scientists

